SEQUENCE LISTING

```
<110> Yan et al.
<120> SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY
<130> 29915/00281F
<140> To be assigned
<141> 2004-03-16
<150> 09/908,943
<151> 2001-07-19
<150> 60/219,795
<151> 2000-07-19
<160> 197
<170> PatentIn Ver. 2.0
<210> 1
<211> 2070
<212> DNA
<213> Homo sapiens
<400> 1
atggcccaag ccctgccctg gctcctgctg tggatggcg cgggagtgct gcctgcccac 60
ggcacccage aeggcateeg getgeeeetg egeageggee tggggggege eeeeetgggg 120
gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
gtgggcagcc ccccgcagac gctcaacatc ctggtggata caggcagcag taactttgca 300
gtgggtgctg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 360
taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
ctgggcaccg acctggtaag catcccccat ggccccaacg tcactgtgcg tgccaacatt 480
getgecatea etgaateaga eaagttette ateaaegget eeaaetggga aggeateetg 540
gggctggcct atgctgagat tgccaggcct gacgactccc tggagccttt ctttgactct 600
ctggtaaagc agacccacgt toccaacctc ttctccctgc acctttgtgg tgctggcttc 660
cccctcaacc agtctgaagt gctggcctct gtcggaggga gcatgatcat tggaggtatc 720
gaccactege tgtacacagg cagtetetgg tatacaceca teeggeggga gtggtattat 780
gaggtcatca ttgtgcgggt ggagatcaat ggacaggatc tgaaaatgga ctgcaaggag 840
tacaactatg acaagagcat tgtggacagt ggcaccacca accttcgttt gcccaaqaaa 900
gtgtttgaag ctgcagtcaa atccatcaag gcagcctcct ccacggagaa gttccctgat 960
ggtttctggc taggagagca gctggtgtgc tggcaagcag gcaccacccc ttggaacatt 1020
ttcccagtca tctcactcta cctaatgggt gaggttacca accagtcctt ccgcatcacc 1080
atcetteege ageaatacet geggeeagtg gaagatgtgg ceaegteeea agacgaetgt 1140
tacaagtttq ccatctcaca qtcatccacq qqcactqtta tqqqaqctqt tatcatqqaq 1200
ggcttctacg ttgtctttga tcgggcccga aaacgaattg gctttgctgt cagcgcttgc 1260
catgtgcacg atgagttcag gacggcagcg gtggaaggcc cttttgtcac cttggacatg 1320
gaagactgtg gctacaacat tccacagaca gatgagtcaa ccctcatgac catagcctat 1380
gtcatggctg ccatctgcgc cctcttcatg ctgccactct gcctcatggt gtgtcagtgg 1440
egetgeetee getgeetgeg ceageageat gatgaetttg etgatqaeat eteeetgetq 1500
aagtgaggag gcccatgggc agaagataga gattcccctg gaccacacct ccgtggttca 1560
ctttggtcac aagtaggaga cacagatggc acctgtggcc agagcacctc aggaccctcc 1620
ccacccacca aatgcctctg ccttgatgga gaaggaaaag gctggcaagg tgggttccag 1680
ggactgtacc tgtaggaaac agaaaagaga agaaagaagc actctqctqq cqqqaatact 1740
cttggtcacc tcaaatttaa gtcgggaaat tctgctgctt gaaacttcag ccctgaacct 1800
gtactggcat cacacgcagg ttaccttggc gtgtgtccct gtggtaccct ggcagagaag 1920
agaccaaget tgttteeetg etggeeaaag teagtaggag aggatgeaca gtttgetatt 1980
tgctttagag acagggactg tataaacaag cctaacattg gtgcaaagat tgcctcttga 2040
attaaaaaaa aaaaaaaaa aaaaaaaaaa
                                                               2070
```

<210> 2

<211> 501

<212> PRT

<213> Homo sapiens

<400> 2

Met Ala Gln Ala Leu Pro Trp Leu Leu Leu Trp Met Gly Ala Gly Val

Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser 20 25 30

Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp 35 40 45

Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
50 55 60

Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr 65 70 75 80

Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
85 90 95

Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr 100 105 110

Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val 115 120 125

Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp 130 135 140

Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile 145 150 155 160

Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp 165 170 175

Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp 180 185 190

Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro 195 200 205

Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln 210 215 220

Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile 225 230 235 240

Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg 245 250 255

Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln 260 265 270

Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val 275 280 285

Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala 290 295 300

```
Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp
                                       315
Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr
                                   330
Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val
Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
                       375
Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
385
Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
                               425
Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
                           440
Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
                       455
Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp
                                  490
Ile Ser Leu Leu Lys
           500
<210> 3
<211> 1977
<212> DNA
<213> Homo sapiens
<400>3
atggcccaag ccctgccctg gctcctgctg tggatggcg cgggagtgct gcctgcccac 60
ggcacccage acggcatecg getgeceetg egeageggee tggggggege ecceetgggg 120
gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
gtgggcagcc ccccgcagac gctcaacatc ctggtggata caggcagcag taactttgca 300
gtgggtgctg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 360
taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
ctgggcaccg acctggtaag catcccccat ggccccaacg tcactgtgcg tgccaacatt 480
gctgccatca ctgaatcaga caagttette atcaacgget ccaactggga aggeateetg 540
gggctggcct atgctgagat tgccaggctt tgtggtgctg gcttccccct caaccagtct 600
gaagtgctgg cctctgtcgg agggagcatg atcattggag gtatcgacca ctcgctgtac 660
acaggcagtc tetggtatac acceateegg egggagtggt attatgaggt gateattgtg 720
```

cgggtggaga tcaatggaca ggatctgaaa atggactgca aggagtacaa ctatgacaag 780 agcattgtgg acagtggcac caccaacctt cgtttgccca agaaagtgtt tgaagctgca 840 gtcaaatcca tcaaggcagc ctcctccacg gagaagttcc ctgatggttt ctggctagga 900 gagcagctgg tgtgctggca agcaggcacc accccttgga acattttccc agtcatctca 960 ctctacctaa tgggtgaggt taccaaccag tccttccgca tcaccatcct tccgcagcaa 1020

```
tacctgcggc cagtggaaga tgtggccacg tcccaagacg actgttacaa gtttgccatc 1080
tcacagtcat ccacgggcac tgttatggga gctgttatca tggagggctt ctacgttgtc 1140
tttgatcggg cccgaaaacg aattggcttt gctgtcagcg cttgccatgt gcacgatgag 1200
ttcaggacgg cagcggtgga aggccctttt gtcaccttgg acatggaaga ctgtggctac 1260
aacattccac agacagatga gtcaaccctc atgaccatag cctatgtcat ggctgccatc 1320
tgcgccetct tcatgctgcc actctgcctc atggtgtgtc agtggcgctg cctccgctgc 1380
ctgcgccage agcatgatga ctttgctgat gacatctccc tgctgaagtg aggaggccca 1440
tgggcagaag atagagattc ccctggacca cacctccgtg gttcactttg gtcacaagta 1500
ggagacacag atggcacctg tggccagagc acctcaggac cctccccacc caccaaatgc 1560
ctctgccttg atggagaagg aaaaggctgg caaggtgggt tccagggact gtacctgtag 1620
gaaacagaaa agagaagaaa gaagcactet getggeggga ataetettgg teaceteaaa 1680
tttaagtcgg gaaattctgc tgcttgaaac ttcagccctg aacctttgtc caccattcct 1740
ttaaattctc caacccaaag tattcttctt ttcttagttt cagaagtact ggcatcacac 1800
gcaggttacc ttggcgtgtg tccctgtggt accctggcag agaagagacc aagcttgttt 1860
ccctgctggc caaagtcagt aggagaggat gcacagtttg ctatttgctt tagagacagg 1920
gactgtataa acaagcctaa cattggtgca aagattgcct cttgaaaaaa aaaaaaa
<210> 4
<211> 476
<212> PRT
<213> Homo sapiens
<400> 4
Met Ala Gln Ala Leu Pro Trp Leu Leu Trp Met Gly Ala Gly Val
Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
                            120
Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
                    150
                                        155
Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
                165
                                    170
Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Leu Cys Gly
Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
```

210 215 220 Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val 230 235 Arg Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser 275 280 Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser 315 Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile 330 Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val 360 Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu 390. Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu 405 410 Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp Ile Ser Leu Leu Lys 470 <210> 5 <211> 14 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic

Lys Val Glu Ala Asn Tyr Glu Val Glu Gly Glu Arg Lys Lys

peptide sequence

<400> 5

10 <210> б <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 6 Lys Val Glu Ala Asn Tyr Glu Val Glu Gly Glu Arg Cys Lys 5 10 <210> 7 <211> 14 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: synthetic peptide sequence <400> 7 Lys Val Glu Ala Asn Tyr Ala Val Glu Gly Glu Arg Lys Lys 5 <210> 8 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 8 Lys Val Glu Ala Asn Tyr Ala Val Glu Gly Glu Arg Cys Lys 5 10 <210> 9 <211> 8 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 9 Glu Ala Asn Tyr Glu Val Glu Phe 5

<210> 10 <211> 8 <212> PRT

<213> Artificial Sequence

```
<220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 10
 Gly Val Leu Leu Ala Ala Gly Trp
 <210> 11
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 11
 Ile Ile Lys Met Asp Asn Phe Gly
 <210> 12
 <211> 10
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <400> 12
 Asp Ser Ser Asn Leu Glu Met Thr His Ala
 <210> 13
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (7)
 <223> Xaa=cysteic acid
 <400> 13
 Thr His Gly Phe Gln Leu Xaa His
 <210> 14
 <211> 8
 <212> PRT
 <213> Artificial Sequence
. <220>
 <223> Description of Artificial Sequence: synthetic
```

-8peptide sequence <400> 14 Cys Tyr Thr His Ser Phe Ser Pro <210> 15 <211> 8 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <220> <221> SITE <222> (4) <223> Xaa= any amino acid <220> <221> SITE <222> (7) <223> Xaa= any amino acid <400> 15 Ser Thr Phe Xaa Gly Ser Xaa Gly 1 <210> 16 <211> 8 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: synthetic peptide sequence <220> <221> SITE <222> (1) <223> Xaa= any amino acid <220> <221> SITE <222> (4)..(7) <223> Xaa= any amino acid <400> 16 Xaa Phe Ala Xaa Xaa Xaa Asn 1 5 <210> 17 <211> 8

<223> Description of Artificial Sequence: synthetic

<212> PRT

<220>

<213> Artificial Sequence

peptide sequence

```
<220>
<221> SITE
<222> (1)..(2)
<223> Xaa=any amino acid
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
Xaa Xaa Gln Xaa Xaa Xaa Ser
 1
<210> 18
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)..(2)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
<400> 18
Xaa Xaa Glu Xaa Xaa Xaa Glu
<210> 19
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 19
Ser Glu Val Asn Leu Asp Ala Glu Phe Arg
<210> 20
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 20
```

```
Ser Glu Val Lys Met Asp Ala Glu Phe Arg
<210> 21
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> MOD RES
<222> (5)
<223> Nle
Ser Glu Val Asn Xaa Asp Ala Glu Phe Arg
                 5
<210> 22
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 22
Gly Ser Glu Ser Met Asp Ser Gly Ile Ser Leu Asp Asn Lys Trp
<210> 23
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 23
Trp Lys Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Lys
                  5
Lys
<210> 24
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
```

```
<400> 24
Ala Asn Leu Ser Thr Phe Ala Gln Pro Arg Arg
                 5
<210> 25
<211> 20
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys Leu Val
                                      10
Phe Phe Ala Glu
<210> 26
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 26
Leu Thr Gly Lys Thr Ile Thr Leu Glu Val Glu Pro Ser Asp Thr Ile
                  5
                                      10
<210> 27
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (19)
<223> Xaa = cysteic acid
<400> 27
Phe Val Asn Gln His Leu Xaa Gly Ser His Leu Val Glu Ala Leu Tyr
Leu Val Xaa Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Ala
             20
                                  25
```

```
<210> 28
<211> 21
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
<220>
<221> SITE
<222> (6)
<223> Xaa=cysteic acid
<220>
<221> SITE
<222> (7)
<223> Xaa=cysteic acid
<220>
<221> SITE
<222> (11)
<223> Xaa=cysteic acid
<220>
<221> SITE
<222> (20)
<223> Xaa=cysteic acid
<400> 28
Gly Ile Val Glu Gln Xaa Xaa Ala Ser Val Xaa Ser Leu Tyr Gln Leu
                                      10
Glu Asn Tyr Xaa Asn
             20
<210> 29
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 29
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
 1
                  5
                                      10
                                                           15
Leu His Ala Leu Gly Gly Cys
             20
<210> 30
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
```

```
<400> 30
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
                 5
Leu His Ala Leu Gly Gly Cys
<210> 31
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 31
Leu Val Asn Met Ala Glu Gly Asp
<210> 32
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 32
Arg Gly Ser Met Ala Gly Val Leu
<210> 33
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 33
Gly Thr Gln His Gly Ile Arg Leu
<210> 34
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 34
Ser Ser Asn Phe Ala Val Gly Ala
                  5
```

```
<210> 35
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 35
Gly Leu Ala Tyr Ala Glu Ile Ala
                  5
<210>. 36
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 36
His Leu Cys Gly Ser His Leu Val
                5
<210> 37
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 37
Cys Gly Glu Arg Gly Phe Phe Tyr
<210> 38
<211> 7
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 38
Gly Val Leu Leu Ser Arg Lys
<210> 39
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
```

```
peptide sequence
<400> 39
Val Gly Ser Gly Val Leu Leu
<210> 40
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 40
Val Gly Ser Gly Val
<210> 41
<211> 12
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (9)
<223> Xaa= cysteic acid
Lys Val Glu Ala Leu Tyr Leu Val Xaa Gly Glu Arg
                 5
<210> 42
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 42
Trp Arg Arg Val Glu Ala Leu Tyr Leu Val Glu Gly Glu Arg Lys
<210> 43
<211> 14
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
```

```
<400> 43
Lys Val Glu Ala Asn Tyr Leu Val Glu Glu Glu Arg Lys Lys
                 5
<210> 44
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 44
Met Leu Leu Leu
 1
<210> 45
<211> 6
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 45
Asp Ala Ala His Pro Gly
<210> 46
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 46
Lys Val Glu Ala Asn Tyr Asp Val Glu Gly Glu Arg Lys Lys
 1
                 5
                                      10
<210> 47
<211> 14
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 47
Lys Val Glu Ala Asn Leu Ala Val Glu Gly Glu Arg Lys Lys
                 5
<210> 48
<211> 14
```

```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 48
Lys Val Glu Ala Leu Tyr Ala Val Glu Gly Glu Arg Lys Lys
<210> 49
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa = E, G, I, D, T, cysteic acid or S
<400> 49
Xaa Ala Asn Tyr Glu Val Glu Phe
  1
                  5
<210> 50
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (2)
<223> Xaa= A, V, I, S, H, Y, T or F
Glu Xaa Asn Tyr Glu Val Glu Phe
<210> 51
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q, or E
```

```
<400> 51
Glu Ala Xaa Tyr Glu Val Glu Phe
                 5
<210> 52
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<400> 52
Glu Ala Asn Xaa Glu Val Glu Phe
<210> 53
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<400> 53
Glu Ala Asn Tyr Xaa Val Glu Phe
 1
<210> 54
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<400> 54
Glu Ala Asn Tyr Glu Xaa Glu Phe
 1
                  5
```

<210> 55

```
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 55
Glu Ala Asn Tyr Glu Val Xaa Phe
 1
<210> 56
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N, S or E
<400> 56
Glu Ala Asn Tyr Glu Val Glu Xaa
<210> 57
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= E, G, I, D, T, cyeteic acid or S
<400> 57
Xaa Val Leu Leu Ala Ala Gly Trp
 1
<210> 58
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
```

```
peptide sequence
<220>
<221> SITE
<222> (2)
<223> Xaa= A, V, I, S, H, Y, T or F
<400> 58
Gly Xaa Leu Leu Ala Ala Gly Trp
<210> 59
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q or E
Gly Val Xaa Leu Ala Ala Gly Trp
<210> 60
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<400> 60
Gly Val Leu Xaa Ala Ala Gly Trp
<210> 61
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
```

```
<400> 61
Gly Val Leu Leu Xaa Ala Gly Trp
<210> 62
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<400> 62
Gly Val Leu Leu Ala Xaa Gly Trp
<210> 63
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 63
Gly Val Leu Leu Ala Ala Xaa Trp
<210> 64
<211> 8
<212> PRT
<213> Artificial Sequence
.<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 64
Gly Val Leu Leu Ala Ala Gly Xaa
```

<210> 65

```
<211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (1)
 <223> Xaa= E, G, I, D, T, cysteic acid or S
 <400> 65
 Xaa Ile Lys Met Asp Asn Phe Gly
   1
 <210> 66
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
    peptide sequence
 <220>
 <221> SITE
 <222> (2)
 <223> Xaa= A, V, I, S, H, Y, T or F
 <400> 66
 Ile Xaa Lys Met Asp Asn Phe Gly
 <210> 67
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
. <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (3)
 <223> Xaa= N, L, K, S, G, T, D, A, Q or E
 <400> 67
 Ile Ile Xaa Met Asp Asn Phe Gly
 <210> 68
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
```

```
peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<400> 68
Ile Ile Lys Xaa Asp Asn Phe Gly
<210> 69
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<400> 69
Ile Ile Lys Met Xaa Asn Phe Gly
                 5
<210> 70
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N,T, L, F or S
<400> 70
Ile Ile Lys Met Asp Xaa Phe Gly
 1
<210> 71
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (7)
```

```
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 71
Ile Ile Lys Met Asp Asn Xaa Gly
<210> 72
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 72
Ile Ile Lys Met Asp Asn Phe Xaa
<210> 73
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= E, G, I, D, T, cysteic acid or S
<400> 73
Xaa Ser Ser Asn Leu Glu Met Thr His Ala
                  5
<210> 74
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (2)
<223> Xaa= A, V, I, S, H, Y, T or F
<400> 74
Asp Xaa Ser Asn Leu Glu Met Thr His Ala
                 5
```

```
<210> 75
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q or E
<400> 75
Asp Ser Xaa Asn Leu Glu Met Thr His Ala
<210> 76
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<400> 76
Asp Ser Ser Xaa Met Thr His Ala
<210> 77
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= E, A, D, M, Q, S or G
<400> 77
Asp Ser Ser Asn Leu Glu Xaa Thr His Ala
<210> 78
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
```

```
<223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (8)
 <223 > Xaa= V, A, N, T, L, F or S
 <400> 78
 Asp Ser Ser Asn Leu Glu Met Xaa His Ala
  1
                   5
 <210> 79
 <211> 9
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <220>
 <221> SITE
 <222> (8)
 <223> Xaa= E, G, F, H, cysteic acid or S
 <400> 79
 Asp Ser Asn Leu Glu Met Thr Xaa Ala
  1
                   5
 <210> 80
 <211> 9
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <220>
 <221> SITE
 <222> (9)
 <223> Xaa= F, W, G, A, H, P, G, N or S
 <400> 80
 Asp Ser Asn Leu Glu Met Thr His Xaa
 <210> 81
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
```

THE P

```
<223> Xaa= E, G, I, D, T, cysteic acid or S
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<400> 81
Xaa His Gly Phe Gln Leu Xaa His
                5
<210> 82
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (2)
<223> Xaa= A, V, I, S, H, Y, T or F
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<400> 82
Thr Xaa Gly Phe Gln Leu Xaa His
 1
<210> 83
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q or E
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<400> 83
Thr His Xaa Phe Gln Leu Xaa His
 1
<210> 84
<211> 8
<212> PRT
```

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<400> 84
Thr His Gly Xaa Gln Leu Xaa His
                  5
<210> 85
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<400> 85
Thr His Gly Phe Xaa Leu Xaa His
                  5
<210> 86
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
```

```
<400> 86
Thr His Gly Phe Gln Xaa Xaa His
 1
<210> 87
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 87
Thr His Gly Phe Gln Leu Xaa His
<210> 88
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (7)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 88
Thr His Gly Phe Gln Leu Xaa Xaa
<210> 89
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= E, G, I, D, T, cysteic acid or S
<400> 89
```

```
Xaa Tyr Thr His Ser Phe Ser Pro
 <210> 90
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (1)
 <223> Xaa= cysteic acid
 <220>
 <221> SITE
 <222> (2)
 <223> Xaa= A, V, I, S, H, Y, T or F
 <400> 90
 Xaa Xaa Thr His Ser Phe Ser Pro
 <210> 91
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
<223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
 <222> (1)
 <223> Xaa= cysteic acid
 <220>
 <221> SITE
 <222> (3)
 <223> Xaa= N, L, K, S, G, T, D, A, Q or E
 <400> 91
 Xaa Tyr Xaa His Ser Phe Ser Pro
 <210> 92
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <220>
 <221> SITE
```

```
<222> (1)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<400> 92
Xaa Tyr Thr Xaa Ser Phe Ser Pro
 1
                 5
<210> 93
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<400> 93
Xaa Tyr Thr His Xaa Phe Ser Pro
 1
<210> 94
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<400> 94
Xaa Tyr Thr His Ser Xaa Ser Pro
 1
                  5
<210> 95
<211> 8
```

```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= cysteic acid
<220>
<221> SITE
<222> (7)
<223> Xaa=E, G, F, H, cysteic acid or S
<400> 95
Xaa Tyr Thr His Ser Phe Xaa Pro
 1
<210> 96
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa=cysteic acid
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 96
Xaa Tyr Thr His Ser Phe Ser Xaa
                  5
<210> 97
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= E, G, I, D, T, cysteic acid or S
<220>
<221> SITE
<222> (7)
```

```
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<400> 97
Xaa Thr Asp Xaa Gly Ser Xaa Gly
<210> 98
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (2)
<223> Xaa=A, V, I, S, H, Y, T or F
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
<400> 98
Ser Xaa Asp Xaa Gly Ser Xaa Gly
<210> 99
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q or E
<220> -
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
```

```
<223> Xaa= any amino acid
<400> 99
Ser Thr Xaa Xaa Gly Ser Xaa Gly
<210> 100
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
<400> 100
Ser Thr Asp Xaa Gly Ser Xaa Gly
<210> 101
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<400> 101
Ser Thr Asp Xaa Xaa Ser Xaa Gly
<210> 102
<211> 8
<212> PRT
```

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<400> 102
Ser Thr Asp Xaa Gly Xaa Xaa Gly
<210> 103
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 103
Ser Thr Asp Xaa Gly Ser Xaa Gly
<210> 104
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (4)
<223> Xaa= any amino acid
```

```
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 104
Ser Thr Asp Xaa Gly Ser Xaa Xaa
<210> 105
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= E, G, I, D, T, cysteic acid or S
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
<400> 105
Xaa Phe Ala Xaa Xaa Xaa Asn
<210> 106
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (2)
<223> Xaa= A, V, I, S, H, Y, T or F
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
```

```
<400> 106
Xaa Xaa Ala Xaa Xaa Xaa Asn
<210> 107
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (3)
<223> Xaa= N, L, K, S, G, T, D, A, Q or E
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
<400> 107
Xaa Phe Xaa Xaa Xaa Xaa Asn
 1
                  5
<210> 108
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)
<223> Xaa= Y, L, M, Nle, F or H
<220>
<221> SITE
<222> (5)..(7)
<223> Xaa= any amino acid
<400> 108
Xaa Phe Ala Xaa Xaa Xaa Asn
 1
                  5
```

```
<210> 109
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)
<223> Xaa = any amino acid
<220>
<221> SITE
<222> (5)
<223> Xaa= E, A, D, M, Q, S or G
<220>
<221> SITE
<222> (6)..(7)
<223> Xaa= any amino acid
<400> 109
Xaa Phe Ala Xaa Xaa Xaa Asn
 1
                  5
<210> 110
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)..(5)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (6)
<223> Xaa= V, A, N, T, L, F or S
<220>
<221> SITE
<222> (7)
<223> Xaa= any amino acid
```

```
<400> 110
Xaa Phe Ala Xaa Xaa Xaa Asn
 1
<210> 111
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)..(6)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 111
Xaa Phe Ala Xaa Xaa Xaa Asn
 1
                 5
<210> 112
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)..(7)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (8)
<223> Xaa= F, W, G, A, H, P, G, N or S
<400> 112
Xaa Phe Ala Xaa Xaa Xaa Xaa
 1
```

```
<210> 113
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 113
Glu Val. Asn Leu Asp Ala Glu Phe Arg
  1
<210> 114
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 114
Asp Tyr Lys Asp Asp Asp Lys
<210> 115
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
Ala Cys Gly Ser Glu Ser Met Asp Ser Gly Ile Ser Leu Asp Asn Lys
Trp
<210> 116
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 116
Trp Lys Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Lys
Lys
```

```
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 117
Ala Asn Leu Ser Thr Phe Ala Gln Pro Arg Arg
                  5
<210> 118
<211> 22
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 118
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
Leu His Leu Gly Gly Cys
             20
<210> 119
<211> 22
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 119
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
                                      10
Leu His Leu Gly Gly Cys
             20
<210> 120
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 120
Lys Thr Ile Thr Leu Glu Val Glu Pro Ser
                  5
<210> 121
<211> 12
```

```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (9)
<223> Xaa= cysteic acid
<400> 121
Val Glu Ala Leu Tyr Leu Val Cys Xaa Gly Glu Arg
                  5
<210> 122
<211> 11
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 122
Val Glu Ala Leu Tyr Leu Val Glu Gly Glu Arg
<210> 123
<211> 363
<212> PRT
<213> Homo sapiens
<220>
<223> galactosyltransferase
<400> 123
Met Ala Ser Lys Ser Trp Leu Asn Phe Leu Thr Phe Leu Cys Gly Ser
                                      10
Ala Ile Gly Phe Leu Leu Cys Ser Gln Leu Phe Ser Ile Leu Leu Gly
             20
Glu Lys Val Asp Thr Gln Pro Asn Val Leu His Asn Asp Pro His Ala
Arg His Ser Asp Asp Asn Gly Gln Asn His Leu Glu Gly Gln Met Asn
Phe Asn Ala Asp Ser Ser Gln His Lys Asp Glu Asn Thr Asp Ile Ala
                     70
Glu Asn Leu Tyr Gln Lys Val Arg Ile Leu Cys Trp Val Met Thr Gly
Pro Gln Asn Leu Glu Lys Lys Ala Lys His Val Lys Ala Thr Trp Ala
Gln Arg Cys Asn Lys Val Leu Phe Met Ser Ser Glu Glu Asn Lys Asp
        115
                            120
                                                 125
```

PhePro 130AlaValGlyLeuLysThrLysGluGlyArgAspGlnLeuTyrTrp 145LysThrIleLysAlaPheGlnTyrValHisGluHisTyrLeuGluAspAlaAspTrpPheLeuLysAlaAspAspThrTyrValIleAspAsnLeuArgTrpLeuLeuSerLysTyrAspProGluGluProIleTyrPheGlyArgArgPheLysProTyrValLysGluLysArgPheValGlyAlaGlyTyrValLeuLysGluAlaLeuLysArgPheVal

Asp Ala Phe Lys Thr Asp Lys Cys Thr His Ser Ser Ser Ile Glu Asp 225 230 235 240

Leu Ala Leu Gly Arg Cys Met Glu Ile Met Asn Val Glu Ala Gly Asp 245 250 255

Ser Arg Asp Thr Ile Gly Lys Glu Thr Phe His Pro Phe Val Pro Glu 260 265 270

His His Leu Ile Lys Gly Tyr Leu Pro Arg Thr Phe Trp Tyr Trp Asn 275 280 285

Tyr Asn Tyr Tyr Pro Pro Val Glu Gly Pro Gly Cys Cys Ser Asp Leu 290 295 300

Ala Val Ser Phe His Tyr Val Asp Ser Thr Thr Met Tyr Glu Leu Glu 305 310 315 320

Tyr Leu Val Tyr His Leu Arg Pro Tyr Gly Tyr Leu Tyr Arg Tyr Gln 325 330 335

Pro Thr Leu Pro Glu Arg Ile Leu Lys Glu Ile Ser Gln Ala Asn Lys 340 345 350

Asn Glu Asp Thr Lys Val Lys Leu Gly Asn Pro 355 360

<210> 124

<211> 405

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens sialylytransferase 1

<400> 124

Ile His Thr Asn Leu Lys Lys Lys Phe Ser Cys Cys Val Leu Val Phe 1 5 10 15

Leu Leu Phe Ala Val Ile Cys Val Trp Lys Glu Lys Lys Gly Ser
20 25 30

Tyr Tyr Asp Ser Phe Lys Leu Gln Thr Lys Glu Phe Gln Val Leu Lys

35 40 45

Ser Leu Gly Lys Leu Ala Met Gly Ser Asp Ser Gln Ser Val Ser Ser 55 Ser Ser Thr Gln Asp Pro His Arg Gly Arg Gln Thr Leu Gly Ser Leu Arg Gly Leu Ala Lys Ala Lys Pro Glu Ala Ser Phe Gln Val Trp Asn Lys Asp Ser Ser Ser Lys Asn Leu Ile Pro Arg Leu Gln Lys Ile Trp 100 105 Lys Asn Tyr Leu Ser Met Asn Lys Tyr Lys Val Ser Tyr Lys Gly Pro Gly Pro Gly Ile Lys Phe Ser Ala Glu Ala Leu Arg Cys His Leu Arg Asp His Val Asn Val Ser Met Val Glu Val Thr Asp Phe Pro Phe Asn 155 Thr Ser Glu Trp Glu Gly Tyr Leu Pro Lys Glu Ser Ile Arg Thr Lys Ala Gly Pro Trp Gly Arg Cys Ala Val Val Ser Ser Ala Gly Ser Leu 180 185 Lys Ser Ser Gln Leu Gly Arg Glu Ile Asp Asp His Asp Ala Val Leu Arg Phe Asn Gly Ala Pro Thr Ala Asn Phe Gln Gln Asp Val Gly Thr 215 Lys Thr Thr Ile Arg Leu Met Asn Ser Gln Leu Val Thr Thr Glu Lys 230 Arg Phe Leu Lys Asp Ser Leu Tyr Asn Glu Gly Ile Leu Ile Val Trp Asp Pro Ser Val Tyr His Ser Asp Ile Pro Lys Trp Tyr Gln Asn Pro Asp Tyr Asn Phe Phe Asn Asn Tyr Lys Thr Tyr Arg Lys Leu His Pro 280 Asn Gln Pro Phe Tyr Ile Leu Lys Pro Gln Met Pro Trp Glu Leu Trp Asp Ile Leu Gln Glu Ile Ser Pro Glu Glu Ile Gln Pro Asn Pro Pro 315 Ser Ser Gly Met Leu Gly Ile Ile Ile Met Met Thr Leu Cys Asp Gln 330 Val Asp Ile Tyr Glu Phe Leu Pro Ser Lys Arg Lys Thr Asp Val Cys 345 Tyr Tyr Tyr Gln Lys Phe Phe Asp Ser Ala Cys Thr Met Gly Ala Tyr 360 His Pro Leu Leu Tyr Glu Lys Asn Leu Val Lys His Leu Asn Gln Gly

370 375 380

Thr Asp Glu Asp Ile Tyr Leu Leu Gly Lys Ala Thr Leu Pro Gly Phe 385 390 395 400

Arg Thr Ile His Cys 405

<210> 125

<211> 518

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens aspartyl protease 1

<400> 125

Met Gly Ala Leu Ala Arg Ala Leu Leu Pro Leu Leu Ala Gln Trp
1 5 10 15

Leu Leu Arg Ala Ala Pro Glu Leu Ala Pro Ala Pro Phe Thr Leu Pro 20 25 30

Leu Arg Val Ala Ala Ala Thr Asn Arg Val Val Ala Pro Thr Pro Gly
35 40 45

Pro Gly Thr Pro Ala Glu Arg His Ala Asp Gly Leu Ala Leu 50 55 60

Glu Pro Ala Leu Ala Ser Pro Ala Gly Ala Ala Asn Phe Leu Ala Met 65 70 75 80

Val Asp Asn Leu Gln Gly Asp Ser Gly Arg Gly Tyr Tyr Leu Glu Met 85 90 95

Leu Ile Gly Thr Pro Pro Gln Lys Leu Gln Ile Leu Val Asp Thr Gly
100 105 1.10

Ser Ser Asn Phe Ala Val Ala Gly Thr Pro His Ser Tyr Ile Asp Thr 115 120 125

Tyr Phe Asp Thr Glu Arg Ser Ser Thr Tyr Arg Ser Lys Gly Phe Asp 130 135 140

Val Thr Val Lys Tyr Thr Gln Gly Ser Trp Thr Gly Phe Val Gly Glu 145 150 155 160

Asp Leu Val Thr Ile Pro Lys Gly Phe Asn Thr Ser Phe Leu Val Asn 165 170 175

Ile Ala Thr Ile Phe Glu Ser Glu Asn Phe Phe Leu Pro Gly Ile Lys 180 185 190

Trp Asn Gly Ile Leu Gly Leu Ala Tyr Ala Thr Leu Ala Lys Pro Ser 195 200 205

Ser Ser Leu Glu Thr Phe Phe Asp Ser Leu Val Thr Gln Ala Asn Ile 210 215 220

Pro Asn Val Phe Ser Met Gln Met Cys Gly Ala Gly Leu Pro Val Ala 225 230 235 240 Gly Ser Gly Thr Asn Gly Gly Ser Leu Val Leu Gly Gly Ile Glu Pro 245 250 255

Ser Leu Tyr Lys Gly Asp Ile Trp Tyr Thr Pro Ile Lys Glu Glu Trp
260 265 270

Tyr Tyr Gln Ile Glu Ile Leu Lys Leu Glu Ile Gly Gly Gln Ser Leu 275 280 285

Asn Leu Asp Cys Arg Glu Tyr Asn Ala Asp Lys Ala Ile Val Asp Ser 290 295 300

Gly Thr Thr Leu Leu Arg Leu Pro Gln Lys Val Phe Asp Ala Val 305 310 315 320

Glu Ala Val Ala Arg Ala Ser Leu Ile Pro Glu Phe Ser Asp Gly Phe 325 330 335

Trp Thr Gly Ser Gln Leu Ala Cys Trp Thr Asn Ser Glu Thr Pro Trp 340 345 350

Ser Tyr Phe Pro Lys Ile Ser Ile Tyr Leu Arg Asp Glu Asn Ser Ser 355 360 365

Arg Ser Phe Arg Ile Thr Ile Leu Pro Gln Leu Tyr Ile Gln Pro Met 370 380

Met Gly Ala Gly Leu Asn Tyr Glu Cys Tyr Arg Phe Gly Ile Ser Pro 385 390 395 4.00

Ser Thr Asn Ala Leu Val Ile Gly Ala Thr Val Met Glu Gly Phe Tyr 405 410 415

Val Ile Phe Asp Arg Ala Gln Lys Arg Val Gly Phe Ala Ala Ser Pro 420 425 430

Cys Ala Glu Ile Ala Gly Ala Ala Val Ser Glu Ile Ser Gly Pro Phe 435 440 445

Ser Thr Glu Asp Val Ala Ser Asn Cys Val Pro Ala Gln Ser Leu Ser 450 460

Glu Pro Ile Leu Trp Ile Val Ser Tyr Ala Leu Met Ser Val Cys Gly
465 470 475 480

Ala Ile Leu Leu Val Leu Ile Val Leu Leu Leu Pro Phe Arg Cys 485 490 495

Gln Arg Arg Pro Arg Asp Pro Glu Val Val Asn Asp Glu Ser Ser Leu 500 505 510

Val Arg His Arg Trp Lys 515

<210> 126

<211> 255

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens syntaxin 6

<400> 126

Met Ser Met Glu Asp Pro Phe Phe Val Val Lys Gly Glu Val Gln Lys

1 10 15

Ala Val Asn Thr Ala Gln Gly Leu Phe Gln Arg Trp Thr Glu Leu Leu 20 25 30

Gln Asp Pro Ser Thr Ala Thr Arg Glu Glu Ile Asp Trp Thr Thr Asn 35 40 45

Glu Leu Arg Asn Asn Leu Arg Ser Ile Glu Trp Asp Leu Glu Asp Leu 50 60

Asp Glu Thr Ile Ser Ile Val Glu Ala Asn Pro Arg Lys Phe Asn Leu 65 70 75 80

Asp Ala Thr Glu Leu Ser Ile Arg Lys Ala Phe Ile Thr Ser Thr Arg 85 90 95

Gln Val Val Arg Asp Met Lys Asp Gln Met Ser Thr Ser Ser Val Gln 100 105 110

Ala Leu Ala Glu Arg Lys Asn Arg Gln Ala Leu Leu Gly Asp Ser Gly
115 120 125

Ser Gln Asn Trp Ser Thr Gly Thr Thr Asp Lys Tyr Gly Arg Leu Asp 130 135 140

Arg Glu Leu Gln Arg Ala Asn Ser His Phe Ile Glu Glu Gln Gln Ala 145 150 155 160

Gln Gln Leu Ile Val Glu Gln Asp Glu Gln Leu Glu Leu Val. 165 170 175

Ser Gly Ser Ile Gly Val Leu Lys Asn Met Ser Gln Arg Ile Gly Gly 180 185 190

Glu Leu Glu Glu Gln Ala Val Met Leu Glu Asp Phe Ser His Glu Leu 195 200 205

Glu Ser Thr Gln Ser Arg Leu Asp Asn Val Met Lys Lys Leu Ala Lys 210 220

Val Ser His Met Thr Ser Asp Arg Gln Trp Cys Ala Ile Ala Ile 225 230 235 240

Leu Phe Ala Val Leu Leu Val Val Leu Ile Leu Phe Leu Val Leu 245 250 250

<210> 127

<211> 1728

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: nucleic acid encoding recombinant fusion protein

<400> 127

atgctgctgc tgctgctgct gctgggcctg aggctacagc tctccctggg catcatccca 60 gttgaggagg agaacccgga cttctggaac cgcgaggcag ccgaggcct gggtgccgcc 120 aagaagctgc agcctgcaca gacagccgcc aagaacctca tcatcttcct gggcgatggg 180

```
atgggggtgt ctacggtgac agctgccagg atcctaaaag ggcagaagaa ggacaaactg 240
gggcctgaga tacccctggc catggaccgc ttcccatatg tggctctgtc caagacatac 300
aatgtagaca aacatgtgcc agacagtgga gccacagcca cggcctacct gtgcggggtc 360
aagggcaact tccagaccat tggcttgagt gcagccgccc gctttaacca gtgcaacacg 420
acacgcggca acgaggtcat ctccgtgatg aatcgggcca agaaagcagg gaagtcagtg 480
ggagtggtaa ccaccacacg agtgcagcac gcctcgccag ccggcaccta cgcccacacg 540
gtgaaccgca actggtactc ggacgccgac gtgcctgcct cggcccgcca ggaqqqqtqc 600
caggacateg ctaegeaget cateteeaac atggacattg aegtgateet aggtggagge 660
cgaaagtaca tgtttcccat gggaacccca gaccctgagt acccagatga ctacagccaa 720
ggtgggacca ggctggacgg gaagaatctg gtgcaggaat ggctggcgaa gcgccagggt 780
gcccggtatg tgtggaaccg cactgagete atgcaggett ccctggacce gtctgtgacc 840
catctcatgg gtctctttga gcctggagac atgaaatacg agatccaccq agactccaca 900
ctggacccct ccctgatgga gatgacagag gctgccctgc gcctgctgag caggaacccc 960
egeggettet teetettegt ggagggtggt egeategace atggteatea tgaaageagg 1020
gcttaccggg cactgactga gacgatcatg ttcgacgacg ccattgagag ggcgggccag 1080
ctcaccageg aggaggacac getgagecte gteactgeeg accaeteeca eqtettetec 1140
ttcggagget acccetgeg agggagetee atettcggge tggeceetgg caaggeeegg 1200
gacaggaagg cetacaeggt cetectatae ggaaaeggte caggetatgt geteaaggae 1260
ggcgcccggc cggatgttac cgagagcgag agcgggagcc ccgagtatcg gcagcagtca 1320
gcagtgcccc tggacgaaga gaccacgca ggcgaggacg tggcggtgtt cgcgcgcggc 1380
ccgcaggcgc acctggttca cggcgtgcag gagcagacct tcatagcgca cgtcatggcc 1440
ttegeegeet geetggagee etacaeegee tgegaeetgg egeeeeege eggeaeeaee 1500
gacgccgcgc acccaggtaa ctatgaagtt gaattccgaa gagcactcta cgtagagggt 1560
gaaagaggat tettetacae teeaaaggea etetaceteg tagagggtga aagaggatte 1620
ttctacacta gtctcatgac catagcctat gtcatggctg ccatctgcgc cctcttcatg 1680
ctgccactct gcctcatggt ggactacaag gatgatgatg acaagtag
<210> 128
<211> 575
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: recombinant
      fusion protein sequence
<400> 128
Met Leu Leu Leu Leu Leu Gly Leu Arg Leu Gln Leu Ser Leu
Gly Ile Ile Pro Val Glu Glu Asn Pro Asp Phe Trp Asn Arg Glu
Ala Ala Glu Ala Leu Gly Ala Ala Lys Lys Leu Gln Pro Ala Gln Thr
Ala Ala Lys Asn Leu Ile Ile Phe Leu Gly Asp Gly Met Gly Val Ser
Thr Val Thr Ala Ala Arg Ile Leu Lys Gly Gln Lys Lys Asp Lys Leu
Gly Pro Glu Ile Pro Leu Ala Met Asp Arg Phe Pro Tyr Val Ala Leu
Ser Lys Thr Tyr Asn Val Asp Lys His Val Pro Asp Ser Gly Ala Thr
Ala Thr Ala Tyr Leu Cys Gly Val Lys Gly Asn Phe Gln Thr Ile Gly
Leu Ser Ala Ala Ala Arg Phe Asn Gln Cys Asn Thr Thr Arg Gly Asn
                       135
                                            140 .
```

Glu Val Ile Ser Val Met Asn Arg Ala Lys Lys Ala Gly Lys Ser Val Gly Val Val Thr Thr Arg Val Gln His Ala Ser Pro Ala Gly Thr 165 170 Tyr Ala His Thr Val Asn Arg Asn Trp Tyr Ser Asp Ala Asp Val Pro Ala Ser Ala Arg Gln Glu Gly Cys Gln Asp Ile Ala Thr Gln Leu Ile Ser Asn Met Asp Ile Asp Val Ile Leu Gly Gly Gly Arg Lys Tyr Met 215 Phe Pro Met Gly Thr Pro Asp Pro Glu Tyr Pro Asp Asp Tyr Ser Gln Gly Gly Thr Arg Leu Asp Gly Lys Asn Leu Val Gln Glu Trp Leu Ala Lys Arg Gln Gly Ala Arg Tyr Val Trp Asn Arg Thr Glu Leu Met Gln 265 Ala Ser Leu Asp Pro Ser Val Thr His Leu Met Gly Leu Phe Glu Pro Gly Asp Met Lys Tyr Glu Ile His Arg Asp Ser Thr Leu Asp Pro Ser 295 Leu Met Glu Met Thr Glu Ala Ala Leu Arg Leu Leu Ser Arg Asn Pro Arg Gly Phe Phe Leu Phe Val Glu Gly Gly Arg Ile Asp His Gly His His Glu Ser Arg Ala Tyr Arg Ala Leu Thr Glu Thr Ile Met Phe Asp 340 345 Asp Ala Ile Glu Arg Ala Gly Gln Leu Thr Ser Glu Glu Asp Thr Leu Ser Leu Val Thr Ala Asp His Ser His Val Phe Ser Phe Gly Gly Tyr 375 Pro Leu Arg Gly Ser Ser Ile Phe Gly Leu Ala Pro Gly Lys Ala Arg Asp Arg Lys Ala Tyr Thr Val Leu Leu Tyr Gly Asn Gly Pro Gly Tyr Val Leu Lys Asp Gly Ala Arg Pro Asp Val Thr Glu Ser Glu Ser Gly 425 Ser Pro Glu Tyr Arg Gln Gln Ser Ala Val Pro Leu Asp Glu Glu Thr His Ala Gly Glu Asp Val Ala Val Phe Ala Arg Gly Pro Gln Ala His Leu Val His Gly Val Gln Glu Gln Thr Phe Ile Ala His Val Met Ala 465 470 475

```
Phe Ala Ala Cys Leu Glu Pro Tyr Thr Ala Cys Asp Leu Ala Pro Pro
Ala Gly Thr Thr Asp Ala Ala His Pro Gly Asn Tyr Glu Val Glu Pro
Arg Arg Ala Leu Tyr Val Glu Gly Glu Arg Gly Phe Phe Tyr Thr Pro
Lys Ala Leu Tyr Leu Val Glu Gly Glu Arg Gly Phe Phe Tyr Thr Ser
Leu Met Thr Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met
Leu Pro Leu Cys Leu Met Val Asp Tyr Lys Asp Asp Asp Lys
<210> 129
<211> 5
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 129
Lys Met Asp Ala Glu
<210> 130
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 130
Gly Arg Arg Gly Ser
 1
<210> 131
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 131
Val Glu Ala Asn Tyr Glu Val Glu Gly Glu
                  5
<210> 132
```

<211> 10

```
<212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <400> 132
 Val Glu Ala Asn Tyr Ala Val Glu Gly Glu
                   5
 <210> 133
 <211> 10
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <400> 133
 Lys Thr Ile Asn Leu Glu Val Glu Pro Ser
  1
             5
 <210> 134
 <211> 10
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
      peptide sequence
 <220>
 <221> MOD_RES
 <222> (5)
 <223> Nle
 <400> 134
 Lys Thr Ile Asn Xaa Glu Val Glu Pro Ser
  1.
                  5
 <210> 135
 <211> 10
 <212> PRT
 <213> Artificial Sequence
 <220>
 <221> MOD_RES
 <222> (5)
 <223> Nle
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 135
Lys Thr Ile Asn Xaa Glu Val Asp Pro Ser
```

5 10 <210> 136 <211> 10 <212> PRT <213> Artificial Sequence <220> <221> MOD RES <222> (5) <223> Nle <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 136 Lys Thr Ile Asn Xaa Asp Val Asp Pro Ser 5 <210> 137 <211> 10 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 137 Lys Thr Ile Ser Leu Asp Val Glu Pro Ser <210> 138 <211> 10 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: synthetic peptide sequence <400> 138 Lys Thr Ile Ser Leu Asp Val Asp Pro Ser <210> 139 <211> 4 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic

peptide sequence

<400> 139

1

Lys Met Asp Ala

```
<210> 140
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 140
Ser Tyr Glu Val
  1
<210> 141
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 141
Ser Glu Val Ser Tyr Glu Val Glu Phe Arg
                  5
<210> 142
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 142
Asn Leu Asp Ala
 1
<210> 143
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 143
Ser Glu Val Ser Tyr Asp Ala Glu Phe Arg
<210> 144
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
```

peptide sequence

```
<400> 144
Ser Glu Val Ser Tyr Glu Ala Glu Phe Arg
<210> 145
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 145
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
Glu Val Ser Tyr Glu Val Glu Phe Arg
             20
<210> 146
<211> 20
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 146
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Val Ser Tyr Glu
                  5
Val Glu Phe Arg
<210> 147
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Lys Thr Glu Glu Ile Ser Glu Val Ser Tyr Glu Val Glu Phe Arg
                  5
<210> 148
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
```

```
<400> 148
Thr Glu Val Ser Tyr Glu Val Glu Phe Arg
                 5
<210> 149
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 149
Ser Glu Val Asp Tyr Glu Val Glu Phe Arg
                  5
<210> 150
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 150
Thr Glu Val Asp Tyr Glu Val Glu Phe Arg
                  5
<210> 151
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 151
Thr Glu Ile Asp Tyr Glu Val Glu Phe Arg
                 5
<210> 152
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 152
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
<210> 153
<211> 10
```

```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 153
Ser Glu Ile Asp Tyr Glu Val Glu Phe Arg
<210> 154
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (11)
<223> Xaa=tryptophan
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
 1
<210> 155
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (16)
<223> Xaa=tryptophan
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 155
Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa
Lys Lys
<210> 156
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (21)
<223> Xaa=tryptophan
<220>
```

```
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 156
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val
                                      10
Glu Phe Arg Xaa Lys Lys
        20
<210> 157
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (26)
<223> Xaa=tryptophan
<400> 157
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
  1
Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 158
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (11)
<223> Xaa=tryptophan
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 158
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 159
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
```

```
<222> (16)
<223> Xaa=tryptophan
<400> 159
Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
Xaa Lys Lys
<210> 160
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (21)
<223> Xaa=tryptophan
<223> Description of Artificial Sequence: synthetic
      peptide
<400> 160
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr
                                      10
Glu Val Glu Phe Arg Xaa Lys Lys
                20
<210> 161
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (26)
<223> Xaa=tryptophan
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 161
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 162
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
```

```
<222> (11)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 162
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 163
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (16)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa
                  5
                                      10
Lys Lys
<210> 164
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (21)
<223> Xaa=oregon green
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 164
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu
                                     10
                                                      15
Val Glu Phe Arg Xaa Lys Lys
             20
<210> 165
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
```

```
<222> (26)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic peptide sequence
<400> 165
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
             20
                                  25
<210> 166
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (11)
<223> Xaa=oregon green
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 166
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
                  5
<210> 167
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (16)
<223> Xaa=oregon green
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 167
Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
Xaa Lys Lys
<210> 168
<211> 23
<212> PRT
<213> Artificial Sequence
```

<220>

```
<221> SITE
<222> (21)
<223> Xaa=oregon green
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 168
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr
                  5
Glu Val Glu Phe Arg Xaa Lys Lys
                20
<210> 169
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (26)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 169
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
                20
<210> 170
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 170
Ser Glu Val Asn Tyr Glu Val Glu Phe Arg
<210> 171
<211> 47
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
<400> 171
gagatetetg aaattagtta tgaagtagaa tteegacatg acteagg
```

```
<210> 172
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
tgagtcatgt cggaattcta cttcataact aatttcagag atctcctc
                                                                   48
<210> 173
<211> 47
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
<400> 173
gagatetetg aaagtagtta tgaagtagaa tteegacatg acteagg
                                                                   47
<210> 174
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
<400> 174
tgagtcatgt cggaattcta cttcataact actttcagag atctcctc
                                                                   48
<210> 175
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
gagatetetg aaattagtta tgaagcagaa tteegacatg acteagg
                                                                   47
<210> 176
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      primer for site-directed mutagenesis of APP
<400> 176
tgagtcatgt cggaattctg cttcataact aatttcagag atctcctc
                                                                   48
<210> 177
<211> 5
<212> PRT
```

. . .

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 177
Val Ser Tyr Glu Val
 1
<210> 178
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 178
Val Ser Tyr Asp Ala
<210> 179
<211> 5
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 179
Ile Ser Tyr Glu Val
 1
<210> 180
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 180
Val Lys Met Asp Ala
<210> 181
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      primer for generating mutant construct named
      MBPC125-SYEV
```

```
<400> 181
gacatetetg aagtgagtta ttaggcagaa tteegacatg acteagg
                                                                    47
<210> 182
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      primer for generating mutant construct named
      MBPC125-SYEV
<400> 182
tgagtcatgt cggaattctg cctaataact cacttcagag atctcctc
                                                                    48
<210> 183
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 183
Lys Lys Ser Tyr Glu Val
 1
                 5
<210> 184
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Val Glu Ala Asn Tyr Glu Val Glu Gly Glu
                  5
<210> 185
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 185
Val Glu Ala Asn Tyr Ala Val Glu Gly Glu
<210> 186
<211> 8
<212> PRT
<213> Artificial Sequence
```

```
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 186
Asp Tyr Lys Asp Asp Asp Lys
<210> 187
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 187
Ser Tyr Glu Ala
 1
<210> 188
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 188
Ser Tyr Ala Val
<210> 189
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 189
Val Ser Tyr Glu Ala
 1
<210> 190
<211> 13
<212> PRT
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic peptide sequence
```

```
<400> 190
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Trp Lys Lys
<210> 191
<211> 23
<212> PRT
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic peptide sequence
<400> 191
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu
Val Glu Phe Arg Trp Lys Lys
            20
<210> 192
<211> 15
<212> PRT
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic peptide sequence
<220>
<221> SITE
<222> (1)..(1)
<223> amino acid at position 1 is biotinylated
<220>
<221> SITE
<222> (14)..(14)
<223> cys at position 14 is derivatized with an oregon green
```

```
<400> 192
Lys Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Lys
<210> 193
<211> 22
<212> PRT
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic peptide sequence
<220>
<221> SITE
<222> (1)..(1)
<223> amino acid at position 1 is biotinylated
<220>
<221> SITE
<222> (21)..(21)
<223> cys at position 21 is derivatized with an oregon green
<400> 193
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu
Val Glu Phe Arg Lys Lys
            20
<210> 194
<211> 6806
<212> DNA
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic DNA sequence
<400> 194
```

ccgacaccat	cgaatggcgc	aaaacctttc	gcggtatggc	atgatagcgc	ccggaagaga	60
gtcaattcag	ggtggtgaat	gtgaaaccag	taacgttata	cgatgtcgca	gagtatgccg	120
gtgtctctta	tcagaccgtt	tcccgcgtgg	tgaaccaggc	cagccacgtt	tctgcgaaaa	180
cgcgggaaaa	agtggaagcg	gcgatggcgg	agctgaatta	cattcccaac	cgcgtggcac	240
aacaactggc	gggcaaacag	tcgttgctga	ttggcgttgc	cacctccagt	ctggccctgc	300
acgcgccgtc	gcaaattgtc	gcggcgatta	aatctcgcgc	cgatcaactg	ggtgccagcg	360
tggtggtgtc	gatggtagaa	cgaagcggcg	tcgaagcctg	taaagcggcg	gtgcacaatc	420
ttctcgcgca	acgcgtcagt	gggctgatca	ttaactatcc	gctggatgac	caggatgcca	480
ttgctgtgga	agctgcctgc	actaatgttc	cggcgttatt	tcttgatgtc	tctgaccaga	540
cacccatcaa	cagtattatt	ttctcccatg	aagacggtac	gcgactgggc	gtggagcatc	600
tggtcgcatt	gggtcaccag	caaatcgcgc	tgttagcggg	cccattaagt	tctgtctcgg	660
cgcgtctgcg	tctggctggc	tggcataaat	atctcactcg	caatcaaatt	cagccgatag	720
cggaacggga	aggcgactgg	agtgccatgt	ccggttttca	acaaaccatg	caaatgctga	780
atgagggcat	cgttcccact	gcgatgctgg	ttgccaacga	tcagatggcg	ctgggcgcaa	840
tgcgcgccat	taccgagtcc	gągctgcgcg	ttggtgcgga	tatctcggta	gtgggatacg	900
acgataccga	agacagctca	tgttatatcc	cgccgttaac	caccatcaaa	caggattttc	960
gcctgctggg	gcaaaccagc	gtggaccgct	tgctgcaact	ctctcagggc	caggcggtga	1020
agggcaatca	gctgttgccc	gtctcactgg	tgaaaagaaa	aaccaccctg	gcgcccaata	1080
cgcaaaccgc	ctctccccgc	gcgttggccg	attcattaat	gcagctggca	cgacaggttt	1140
cccgactgga	aagcgggcag	tgagcgcaac	gcaattaatg	tgagttagct	cactcattag	1200
gcacaattct	catgtttgac	agcttatcat	cgactgcacg	gtgcaccaat	gcttctggcg	1260
tcaggcagcc	atcggaagct	gtggtatggc	tgtgcaggtc	gtaaatcact	gcataattcg	1320
tgtcgctcaa	ggcgcactcc	cgttctggat	aatgttttt	gcgccgacat	cataacggtt	1380
ctggcaaata	ttctgaaatg	agctgttgac	aattaatcat	cggctcgtat	aatgtgtgga	1440
attgtgagcg	gataacaatt	tcacacagga	aacagccagt	ccgtttaggt	gttttcacga	1500
gcacttcacc	aacaaggacc	atagattatg	aaaactgaag	aaggtaaact	ggtaatctgg	1560
attaacggcg	ataaaggcta	taacggtctc	gctgaagtcg	gtaagaaatt	cgagaaagat	1620
accggaatta	aagtcaccgt	tgagcatccg	gataaactgg	aagagaaatt	cccacaggtt	1680
gcggcaactg	gcgatggccc	tgacattatc	ttctgggcac	acgaccgctt	tggtggctac	1740
gctcaatctg	gcctgttggc	tgaaatcacc	ccggacaaag	cgttccagga	caagctgtat	1800
ccgtttacct	gggatgccgt	acgttacaac	ggcaagctga	ttgcttaccc	gatcgctgtt	1860
gaagcgttat	cgctgattta	taacaaagat	ctgctgccga	acccgccaaa	aacctgggaa	1920

			- 09 -			
gagatecegg	cgctggataa	agaactgaaa	gcgaaaggta	agagcgcgct	gatgttcaac	1980
ctgcaagaac	cgtacttcac	ctggccgctg	attgctgctg	acgggggtta	tgcgttcaag	2040
tatgaaaacg	gcaagtacga	cattaaagac	gtgggcgtgg	ataacgctgg	cgcgaaagcg	2100
ggtctgacct	tcctggttga	cctgattaaa	aacaaacaca	tgaatgcaga	caccgattac	2160
tccatcgcag	aagctgcctt	taataaaggc	gaaacagcga	tgaccatcaa	cggcccgtgg	2220
gcatggtcca	acatcgacac	cagcaaagtg	aattatggtg	taacggtact	gccgaccttc	2280
aagggtcaac	catccaaacc	gttcgttggc	gtgctgagcg	caggtattaa	cgccgccagt	2340
ccgaacaaag	agctggcgaa	agagttcctc	gaaaactatc	tgctgactga	tgaaggtctg	2400
gaagcggtta	ataaagacaa	accgctgggt	gccgtagcgc	tgaagtctta	cgaggaagag	2460
ttggcgaaag	atccacgtat	tgccgccacc	atggaaaacg	cccagaaagg	tgaaatcatg	2520
ccgaacatcc	cgcagatgtc	cgctttctgg	tatgccgtgc	gtactgcggt	gatcaacgcc	2580
gccagcggtc	gtcagactgt	cgatgaagcc	ctgaaagacg	cgcagactaa	ttcgagctcg	2640
gtacccggcc	ggggatccat	cgagggtagg	gccgaccgag	gactgaccac	tcgaccaggt	2700 -
tctgggttga	caaatatcaa	gacggaggag	atctctgaag	tgaatctgga	tgcagaattc	2760
cgacatgact	caggatatga	agttcatcat	caaaaattgg	tgttctttgc	agaagatgtg	2820
ggttcaaaca	aaggtgcaat	cattggactc	atggtgggcg	gtgttgtcat	agcgacagtg	2880
atcgtcatca	ccttggtgat	gctgaagaag	aaacagtaca	catccattca	tcatggtgtg	2940 🏄
gtggaggttg	acgccgctgt	caecccagag	gagcgccacc	tgtccaagat	gcagcagaac	3000 .
ggctacgaaa	atccaaccta	caagttcttt	gagcagatgc	agaactagac	ccccgccaca	3060
gcagcctctg	aagttggaca	gcaaaaccat	tgcttcacta	cccatcggtg	tccatttata	3120
gaataatgtg	ggaagaaaca	aacccgtttt	atgatttact	cattatcgcc	ttttgacagc	3180
tgtgctgtaa	cacaagtaga	tgcctgaact	tgaattaatc	cacacatcag	taatgtattc	3240
tatctctctt	tacattttgg	tctctatact	acattattaa	tgggttttgt	gtactgtaaa	3300
gaatttagct	gtatcaaact	agtaatagcc	tgaattcagt	aacctaaccc	tcgatggatc	3360
ctctagagtc	gacctgcagg	caagcttggc	actggccgtc	gttttacaac	gtcgtgactg	3420
ggaaaaccct	ggcgttaccc	aacttaatcg	ccttgcagca	catccccctt	tcgccagctg	3480
gcgtaatagc	gaagaggccc	gcaccgatcg	cccttcccaa	cagttgcgca	gcctgaatgg	3540
cgaatggcag	cttggctgtt	ttggcggatg	agagaagatt	ttcagcctga	tacagattaa	3600
atcagaacgc	agaagcggtc	tgataaaaca	gaatttgcct	ggcggcagta	gcgcggtggt	3660
cccacctgac	cccatgccga	actcagaagt	gaaacgccgt	agcgccgatg	gtagtgtggg	3720
gtctccccat	gcgagagtag	ggaactgcca	ggcatcaaat	aaaacgaaag	gctcagtcga	3780
aagactgggc	ctttcgtttt	atctgttgtt	tgtcggtgaa	agatataatg	agtaggacaa	3840

atccgccggg	agcggatttg	aacgttgcga	agcaacggcc	cggagggtgg	cgggcaggac	3900
gcccgccata	aactgccagg	catcaaatta	agcagaaggc	catcctgacg	gatggccttt	3960
ttgcgtttct	acaaactctt	tttgtttatt	tttctaaata	cattcaaata	tgtatccgct	4020
catgagacaa	taaccctgat	aaatgcttca	ataatattga	aaaaggaaga	gtatgagtat	4080
tcaacatttc	cgtgtcgccc	ttattccctt	ttttgcggca	ttttgccttc	ctgtttttgc	4140
tcacccagaa	acgctggtga	aagtaaaaga	tgctgaagat	cagttgggtg	cacgagtggg	4200
ttacatcgaa	ctggatctca	acagcggtaa	gatccttgag	agttttcgcc	ccgaagaacg	4260
ttttccaatg	atgagcactt	ttaaagttct	gctatgtggc	gcggtattat	cccgtgttga	4320
cgccgggcaa	gagcaactcg	gtcgccgcat	acactattct	cagaatgact	tggttgagta	4380
ctcaccagtc	acagaaaagc	atcttacgga	tggcatgaca	gtaagagaat	tatgcagtgc	4440
tgccataacc	atgagtgata	acactgcggc	caacttactt	ctgacaacga	tcggaggacc	4500
gaaggagcta	accgcttttt	tgcacaacat	gggggatcat	gtaactcgcc	ttgatcgttg	4560
ggaaccggag	ctgaatgaag	ccataccaaa	cgacgagcgt	gacaccacga	tgcctgtagc	4620
aatggcaaca	acgttgcgca	aactattaac	tggcgaacta	cttactctag	cttcccggca	4680
acaattaata	gactggatgg	aggcggataa	agttgcagga	ccacttctgc	gctcggccct	4740
tccggctggc	tggtttattg	ctgataaatc	tggagccggt	gagcgtgggt	ctcgcggtat	4800 %
cattgcagca	ctggggccag	atggtaagcc	ctcccgtatc	gtagttatct	acacgacggg	4860 ⋅ 🖰
gagtcaggca	actatggatg	aacgaaatag	acagatcgct	gagataggtg	cctcactgat	4920
taagcattgg	taactgtcag	accaagttta	ctcatatata	ctttagattg	atttaaaact	4980
tcatttttaa	tttaaaagga	tctaggtgaa	gatccttttt	gataatctca	tgaccaaaat	5040
cccttaacgt	gagttttcgt	tccactgagc	gtcagacccc	gtagaaaaga	tcaaaggatc	5100
ttcttgagat	ccttttttc	tgcgcgtaat	ctgctgcttg	caaacaaaaa	aaccaccgct	5160
accagcggtg	gtttgtttgc	cggatcaaga	gctaccaact	ctttttccga	aggtaactgg	5220
cttcagcaga	gcgcagatac	caaatactgt	ccttctagtg	tagccgtagt	taggccacca	5280
cttcaagaac	tctgtagcac	cgcctacata	cctcgctctg	ctaatcctgt	taccagtggc	5340
tgctgccagt	ggcgataagt	cgtgtcttac	cgggttggac	tcaagacgat	agttaccgga	5400
taaggcgcag	cggtcgggct	gaacgggggg	ttcgtgcaca	cagcccagct	tggagcgaac	5460
gacctacacc	gaactgagat	acctacagcg	tgagctatga	gaaagcgcca	cgcttcccga	5520
agggagaaag	gcggacaggt	atccggtaag	cggcagggtc	ggaacaggag	agcgcacgag	5580
ggagcttcca	gggggaaacg	cctggtatct	ttatagtcct	gtcgggtttc	gccacctctg	5640
acttgagcgt	cgatttttgt	gatgctcgtc	aggggggcgg	agcctatgga	aaaacgccag	5700
caacgcggcc	tttttacggt	tcctggcctt	ttgctggcct	tttgctcaca	tgttctttcc	5760

```
tgcgttatcc cctgattctg tggataaccg tattaccgcc tttgagtgag ctgataccgc
                                                                     5820
tegeegeage egaacgaeeg agegeagega gteagtgage gaggaagegg aagagegeet
                                                                     5880
gatgcggtat tttctcctta cgcatctgtg cggtatttca caccgcatat ggtgcactct
                                                                     5940
cagtacaatc tgctctgatg ccgcatagtt aagccagtat acactccgct atcgctacgt
                                                                     6000
gactgggtca tggctgcgcc ccgacacccg ccaacacccg ctgacgcgcc ctgacgggct
                                                                     6060
tgtctgctcc cggcatccgc ttacagacaa gctgtgaccg tctccqqqaq ctqcatgtgt
                                                                     6120
cagaggtttt caccgtcatc accgaaacgc gcgaggcagc tgcggtaaag ctcatcagcg
                                                                     6180
tggtcgtgaa gcgattcaca gatgtctgcc tgttcatccg cgtccagctc gttgagtttc
                                                                     6240
tccagaagcg ttaatgtctg gcttctgata aagcgggcca tgttaagggc ggttttttcc
                                                                     6300
tgtttggtca cttgatgcct ccgtgtaagg gggaatttct gttcatgggg gtaatgatac
                                                                     6360
cgatgaaacg agagaggatg ctcacgatac gggttactga tgatgaacat gcccggttac
                                                                     6420
tggaacgttg tgagggtaaa caactggcgg tatggatgcg gcgggaccag agaaaaatca
                                                                     6480
ctcagggtca atgccagcgc ttcgttaata cagatgtagg tgttccacag ggtagccagc
                                                                     6540
agcatectge gatgeagate eggaacataa tggtgeaggg egetgaette egegttteea
                                                                     6600
gactttacga aacacggaaa ccgaagacca ttcatgttgt tgctcaggtc gcagacgttt
                                                                     6660
tgcagcagca gtcgcttcac gttcgctcgc gtatcggtga ttcattctgc taaccagtaa
                                                                     6720
ggcaaccccg ccagcctagc cgggtcctca acgacaggag cacgatcatg cgcacccgtg
                                                                     6780
gccaggaccc aacgctgccc gaaatt
                                                                     68.06
```

```
<210> 195
```

<211> 13

<212> PRT

<213> Artificial sequence

<220>

<223> Description of artificial sequence: synthetic peptide sequence

<220>

<221> MOD RES

<222> (1)..(1)

<223> ACETYLATION (MCA)

```
<221> SITE
<222> (11)..(11)
<223> 2,4-dinitrophenyl group after the Lys at position 11
<400> 195
Ser Glu Val Asn Leu Asp Ala Glu Phe Arg Lys Arg Arg
<210> 196
<211> 12
<212> PRT
<213> Artificial sequence
<220>
<223> Description of artificial sequence: synthetic peptide sequence
<220>
<221> SITE
<222> (4)..(4)
<223> amino acid at position 4 has been derivatized with a statine
<400> 196
Ser Glu Val Asn Val Ala Glu Phe Arg Gly Gly Cys
<210> 197
<211> 10
<212> PRT
<213> synthetic peptide sequence
<220>
<221> SITE
<222>
     (4)..(4)
<223> amino acid at position 4 has been derivatized with a statine
<220>
```

<221> SITE

<222> (10)..(10)

<223> amino acid at position 10 has been derivatized with Bodipy FL

<400> 197

Ser Glu Val Asn Val Ala Glu Phe Arg Cys 1 5 10